

HS arch Differential expression HvN

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ID	Image	logFC	From	From	From
			to	to	to
k141_181201_18_-		-7.69	3.73e-09	1.55e-06	1.14e-06
k141_372642_30_-		-7.69	1.92e-06	1.14e-06	1.14e-06
k141_326599_21_CBS_domain		-7.49	5.62e-07	5.35e-07	5.35e-07
k141_93077_13_-		-6.72	1.99e-07	4.15e-07	4.15e-07
k141_326599_17_Universal_stress_protein		-6.38	6.43e-07	5.35e-07	5.35e-07
k141_392539_6_COG0589_Universal_stress_protein_Usp		-6.32	5.21e-07	5.35e-07	5.35e-07
k141_181201_19_Domain_of_unknown_function_DUF302		-6.16	6.60e-05	2.74e-05	2.74e-05
k141_96470_27_CBS_domain		-5.63	6.54e-05	2.74e-05	2.74e-05
k141_92599_11_COG1522_Transcriptional_regulators		-5.59	1.83e-03	5.17e-03	5.17e-03
k141_76157_13_Belongs_to_the_P_II__protein_family		-5.57	1.39e-05	7.22e-05	7.22e-05
k141_92599_5_helix_turn_helix_ASNC_type		-5.12	2.39e-03	6.21e-03	6.21e-03
k141_326599_18_COG0589_Universal_stress_protein_Us		-5.07	2.26e-04	7.85e-04	7.85e-04
k141_440873_29_Belongs_to_the_small_heat_shock_pro		-4.42	1.29e-03	4.13e-03	4.13e-03
k141_447824_6_-		-4.14	1.65e-06	1.14e-06	1.14e-06
k141_92599_3_Lrp_AsnC_ligand_binding_domain		-4.02	1.89e-04	7.16e-04	7.16e-04
k141_440873_33_Stabilizes_TBP_binding_to_an_archae		-3.20	1.86e-03	5.17e-03	5.17e-03
k141_15183_34_-		-2.61	3.97e-03	9.72e-03	9.72e-03

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